

SEQUENCE LISTING

<110> Bandman, Olga
 Tang, Y. Tom
 Hillman, Jennifer L.
 Yue, Henry
 Guegler, Karl J.
 Corley, Neil C.
 Gorgone, Gina
 Azimzai, Yalda
 Lu, Aina

<120> Protein Kinase Homologs

<130> PF-0614 US

<140> To Be Assigned

<141> Herewith

<160> 18

<170> PERL Program

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<212> PRT

<213> Homo sapiens

<220> -

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Leu	Arg	Asp	Leu	Gln	Arg	Leu	Glu	Ala	Met	Glu	Ala	Ala	Thr	Gln	20	25	30	35
Ala	Glu	Asp	Ser	Gly	Leu	Arg	Leu	Asp	Gly	Gly	Ser	Gly	Ser	Thr	40	45	50	55
Ser	Ser	Ser	Gly	Cys	His	Pro	Gly	Gly	Ala	Arg	Ala	Gly	Pro	Ser	60	65	70	75
Pro	Ala	Ser	Ser	Ser	Pro	Ala	Pro	Gly	Gly	Gly	Arg	Ser	Leu	Ser	80	85	90	95
Ala	Gly	Ser	Gln	Thr	Ser	Gly	Phe	Ser	Gly	Ser	Leu	Phe	Ser	Pro	100	105	110	115
Ala	Ser	Cys	Ser	Ile	Leu	Ser	Gly	Ser	Ser	Asn	Gln	Arg	Glu	Thr	120	125	130	135
Gly	Gly	Leu	Leu	Ser	Pro	Ser	Thr	Pro	Phe	Gly	Ala	Ser	Asn	Leu	140	145	150	155
Leu	Val	Asn	Pro	Leu	Glu	Pro	Gln	Asn	Ala	Asp	Lys	Ile	Lys	Ile	160	165	170	175
Lys	Ile	Ala	Asp	Leu	Gly	Asn	Ala	Cys	Trp	Val	His	Lys	His	Phe	180	185	190	195
Thr	Glu	Asp	Ile	Gln	Thr	Arg	Gln	Tyr	Arg	Ala	Val	Glu	Val	Leu	200	205	210	215
Ile	Gly	Ala	Glu	Tyr	Gly	Pro	Pro	Ala	Asp	Ile	Trp	Ser	Thr	Ala	220	225	230	235
Cys	Met	Ala	Phe	Glu	Leu	Ala	Thr	Gly	Asp	Tyr	Leu	Phe	Glu	Pro	240	245	250	255
His	Ser	Gly	Glu	Asp	Tyr	Ser	Arg	Asp	Glu	Asp	His	Ile	Ala	His	250	255	260	265
Ile	Val	Glu	Leu	Leu	Gly	Asp	Ile	Pro	Pro	Ala	Phe	Ala	Leu	Ser	265	270	275	280
Gly	Arg	Tyr	Ser	Arg	Glu	Phe	Phe	Asn	Arg	Arg	Gly	Glu	Leu	Arg	280	285	290	295
His	Ile	His	Asn	Leu	Lys	His	Trp	Gly	Leu	Tyr	Glu	Val	Leu	Met	295	300	305	310

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Glu	Lys	Tyr	Glu	Trp	Pro	Leu	Glu	Gln	Ala	Thr	Gln	Phe	Ser	Ala	
				260					265					270	
Phe	Leu	Leu	Pro	Met	Asn	Glu	Tyr	Ile	Pro	Glu	Lys	Arg	Ala	Ser	
				275					280					285	
Ala	Arg	Asp	Cys	Leu	Gln	His	Pro	Trp	Leu	Gln	Pro				
				290					295						

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<213> Homo sapiens

<220> -
<223> 132750

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Met	Gln	Glu	Ile	Pro	Gln	Glu	Gln	Ile	Lys	Glu	Ile	Lys	Lys	Glu	
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Gln	Leu	Ser	Gly	Ser	Pro	Trp	Ile	Leu	Leu	Arg	Glu	Asn	Glu	Val	
				20					25					30	
Ser	Thr	Leu	Tyr	Lys	Gly	Glu	Tyr	His	Arg	Ala	Pro	Val	Ala	Ile	
				35					40					45	
Lys	Val	Phe	Lys	Lys	Leu	Gln	Ala	Gly	Ser	Ile	Ala	Ile	Val	Arg	
				50					55					60	
Gln	Thr	Phe	Asn	Lys	Glu	Ile	Lys	Thr	Met	Lys	Lys	Phe	Glu	Ser	
				65					70					75	
Pro	Asn	Ile	Leu	Arg	Ile	Phe	Gly	Ile	Cys	Ile	Asp	Glu	Thr	Val	
				80					85					90	
Thr	Pro	Pro	Gln	Phe	Ser	Ile	Val	Met	Glu	Tyr	Cys	Glu	Leu	Gly	
				95					100					105	
Thr	Leu	Arg	Glu	Leu	Leu	Asp	Arg	Glu	Lys	Asp	Leu	Thr	Leu	Gly	
				110					115					120	
Lys	Arg	Met	Val	Leu	Val	Leu	Gly	Ala	Ala	Arg	Gly	Leu	Tyr	Arg	
				125					130					135	
Leu	His	His	Ser	Glu	Ala	Pro	Glu	Leu	His	Gly	Lys	Ile	Arg	Ser	
				140					145					150	
Ser	Asn	Phe	Leu	Val	Thr	Gln	Gly	Tyr	Gln	Val	Lys	Leu	Ala	Gly	
				155					160					165	
Phe	Glu	Leu	Arg	Lys	Thr	Gln	Thr	Ser	Met	Ser	Leu	Gly	Thr	Thr	
				170					175					180	
Arg	Glu	Lys	Thr	Asp	Arg	Val	Lys	Ser	Thr	Ala	Tyr	Leu	Ser	Pro	
				185					190					195	
Gln	Glu	Leu	Glu	Asp	Val	Phe	Tyr	Gln	Tyr	Asp	Val	Lys	Ser	Glu	
				200					205					210	
Ile	Tyr	Ser	Phe	Gly	Ile	Val	Leu	Trp	Glu	Ile	Ala	Thr	Gly	Asp	
				215					220					225	
Ile	Pro	Phe	Gln	Gly	Cys	Asn	Ser	Glu	Lys	Ile	Arg	Lys	Leu	Val	
				230					235					240	
Ala	Val	Lys	Arg	Gln	Gln	Glu	Pro	Leu	Gly	Glu	Asp	Cys	Pro	Ser	
				245					250					255	
Glu	Leu	Arg	Glu	Ile	Ile	Asp	Glu	Cys	Arg	Ala	His	Asp	Pro	Ser	
				260					265					270	
Val	Arg	Pro	Ser	Val	Asp	Glu	Ile	Leu	Lys	Lys	Leu	Ser	Thr	Phe	
				275					280					285	

Ser Lys

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PF-0614 US

<220> -

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Asn Ile Asp Val Cys Glu Asn Cys His Tyr Pro Ile Val Pro Leu
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Asp Gly Lys Gly Thr Leu Leu Ile Arg Asn Gly Ser Glu Val Arg
35 40 45
Asp Pro Leu Val Thr Tyr Glu Gly Ser Asn Pro Pro Ala Ser Pro
50 55 60
Leu Gln Asp Asn Leu Val Ile Ala Leu His Ser Tyr Glu Pro Ser
65 70 75
His Asp Gly Asp Leu Gly Phe Glu Lys Gly Glu Gln Leu Arg Ile
80 85 90
Leu Glu Gln Ser Gly Glu Trp Trp Lys Ala Gln Ser Leu Thr Thr
95 100 105
Gly Gln Glu Gly Phe Ile Pro Phe Asn Phe Val Ala Lys Ala Asn
110 115 120
Ser Leu Glu Pro Glu Ala Asn Leu Met Lys Gln Leu Gln His Gln
125 130 135
Arg Leu Val Arg Leu Tyr Ala Val Val Thr Gln Glu Pro Ile Tyr
140 145 150
Ile Ile Thr Glu Tyr Met Glu Asn Gly Ser Leu Val Asp Phe Leu
155 160 165
Lys Thr Pro Ser Gly Ile Lys Leu Thr Ile Asn Lys Leu Leu Asp
170 175 180
Met Ala Ala Gln Ile Ala Glu Gly Met Ala Phe Ile Glu Glu Arg
185 190 195
Asn Tyr Ile His Arg Asp Leu Arg Ala Ala Asn Ile Leu Val Ser
200 205 210
Asp Thr Leu Ser Cys Lys Ile Ala Asp Phe Gly Leu Ala Arg Leu
215 220 225
Ile Glu Asp Asn Glu Tyr Thr Ala Arg Glu Gly Ala Lys Phe Pro
230 235 240
Ile Lys Trp Thr Ala Pro Glu Ala Ile Asn Tyr Gly Thr Phe Thr
245 250 255
Ile Lys Ser Asp Val Trp Ser Phe Gly Ile Leu Leu Thr Glu Ile
260 265 270
Val Thr His Gly Arg Ile Pro Tyr Pro Gly Met Thr Asn Pro Glu
275 280 285

Val Ile Gln Asn Leu Glu Arg Gly Tyr Arg Met Val Arg Pro Asp
290 295 300
Asn Cys Pro Glu Glu Leu Tyr Gln Leu Met Arg Leu Cys Trp Lys
305 310 315
Glu Arg Pro Glu Asp Arg Pro Thr Phe Asp Tyr Leu Arg Ser Val
320 325 330
Leu Glu Asp Phe Phe Thr Ala Thr Glu Gly Gln Tyr Gln Pro Gln
335 340 345
Pro

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Met Pro Ala Gly Gly Arg Ala Gly Ser Leu Lys Asp Pro Asp Val
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PF-0614 US

Ala	Glu	Leu	Phe	Phe	Lys	Asp	Asp	Pro	Glu	Lys	Leu	Phe	Ser	Asp	
				20					25					30	
Leu	Arg	Glu	Ile	Gly	His	Gly	Ser	Phe	Gly	Ala	Val	Tyr	Phe	Ala	
				35					40					45	
Arg	Asp	Val	Arg	Asn	Ser	Glu	Val	Val	Ala	Ile	Lys	Lys	Met	Ser	
				50					55					60	
Tyr	Ser	Gly	Lys	Gln	Ser	Asn	Glu	Lys	Trp	Gln	Asp	Ile	Ile	Lys	
				65					70					75	
Glu	Val	Arg	Arg	Arg	Arg	Arg	Val	Gly	Arg	Glu	Asp	Glu	Glu	Arg	
				80					85					90	

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Val	Ile	Gly	Lys	Gly	Phe	Glu	Asp	Leu	Met	Thr	Val	Asn	Leu	Ala	
				20					25					30	
Arg	Tyr	Lys	Pro	Thr	Gly	Glu	Tyr	Val	Thr	Val	Arg	Arg	Ile	Asn	
				35					40					45	
Leu	Glu	Ala	Cys	Ser	Asn	Glu	Met	Val	Thr	Phe	Leu	Gln	Gly	Glu	
				50					55					60	
Leu	His	Val	Ser	Lys	Leu	Phe	Asn	His	Pro	Asn	Ile	Val	Pro	Tyr	
				65					70					75	
Arg	Ala	Thr	Phe	Ile	Ala	Asp	Asn	Glu	Leu	Trp	Val	Val	Thr	Ser	
				80					85					90	
Phe	Met	Ala	Tyr	Gly	Ser	Ala	Lys	Asp	Leu	Ile	Cys	Thr	His	Phe	
				95					100					105	
Met	Asp	Gly	Met	Asn	Glu	Leu	Ala	Ile	Ala	Tyr	Ile	Leu	Gln	Gly	
				110					115					120	
Val	Leu	Lys	Ala	Leu	Asp	Tyr	Ile	His	His	Met	Gly	Tyr	Val	His	
				125					130					135	
Arg	Ser	Val	Lys	Ala	Ser	His	Ile	Leu	Ile	Ser	Val	Asp	Gly	Lys	
				140					145					150	
Val	Tyr	Leu	Ser	Gly	Leu	Arg	Thr	Thr	Leu	Ser	Met	Ile	Ser	His	
				155					160					165	
Gly	Gln	Arg	Gln	Arg	Val	Val	His	Asp	Phe	Pro	Lys	Tyr	Ser	Val	
				170					175					180	
Lys	Val	Leu	Pro	Trp	Leu	Ser	Pro	Glu	Val	Leu	Gln	Gln	Asn	Leu	
				185					190					195	
Gln	Gly	Tyr	Asp	Ala	Lys	Ser	Asp	Ile	Tyr	Ser	Val	Gly	Ile	Thr	
				200					205					210	
Ala	Cys	Glu	Leu	Ala	Asn	Gly	His	Val	Pro	Phe	Lys	Asp	Met	Pro	
				215					220					225	
Ala	Thr	Gln	Met	Leu	Leu	Glu	Lys	Leu	Asn	Gly	Thr	Val	Pro	Cys	
				230					235					240	
Leu	Leu	Asp	Thr	Ser	Thr	Ile	Pro	Ala	Glu	Glu	Leu	Thr	Met	Ser	
				245					250					255	
Pro	Ser	Arg	Ser	Val	Ala	Asn	Ser	Gly	Leu	Ser	Asp	Ser	Leu	Thr	
				260					265					270	
Thr	Ser	Thr	Pro	Arg	Pro	Ser	Asn	Gly	Asp	Ser	Pro	Ser	His	Pro	
				275					280					285	
Tyr	His	Arg	Thr	Phe	Ser	Pro	His	Phe	His	His	Phe	Val	Glu	Gln	
				290					295					300	
Cys	Leu	Gln	Arg	Asn	Pro	Asp	Ala	Arg	Tyr	Pro	Cys	Trp	Pro	Gly	
				305					310					315	
Pro	Gly	Leu	Arg	Glu	Ser	Arg	Gly	Cys	Ser	Gly	Gly				
				320					325						

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Met	Glu	Lys	Tyr	Val	Arg	Leu	Gln	Lys	Ile	Gly	Glu	Gly	Ser	Phe
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Gly	Lys	Ala	Ile	Leu	Val	Lys	Ser	Thr	Glu	Asp	Gly	Arg	Gln	Tyr
				20					25					30
Val	Ile	Lys	Glu	Ile	Asn	Ile	Ser	Arg	Met	Ser	Ser	Lys	Glu	Arg
				35					40					45
Glu	Glu	Ser	Arg	Arg	Glu	Val	Ala	Val	Leu	Ala	Asn	Met	Lys	His
				50					55					60
Pro	Asn	Ile	Val	Gln	Tyr	Arg	Glu	Ser	Phe	Glu	Gly	Ile	Leu	Asp
				65					70					75
Trp	Phe	Val	Gln	Ile	Cys	Leu	Ala	Leu	Lys	His	Val	His	Asp	Arg
				80					85					90
Lys	Ile	Leu	His	Arg	Asp	Ile	Lys	Ser	Gln	Asn	Ile	Phe	Leu	Thr
				95					100					105
Lys	Asp	Gly	Thr	Val	Gln	Leu	Gly	Asp	Phe	Gly	Ile	Ala	Arg	Val
				110					115					120
Leu	Asn	Ser	Thr	Val	Glu	Leu	Ala	Arg	Thr	Cys	Ile	Gly	Thr	Pro
				125					130					135
Tyr	Tyr	Leu	Ser	Pro	Glu	Ile	Cys	Glu	Asn	Lys	Pro	Tyr	Asn	Asn
				140					145					150
Lys	Ser	Asp	Ile	Trp	Ala	Leu	Gly	Cys	Val	Leu	Tyr	Glu	Leu	Cys
				155					160					165
Thr	Leu	Lys	His	Ala	Phe	Glu	Ala	Gly	Ser	Met	Lys	Asn	Leu	Val
				170					175					180
Leu	Lys	Ile	Ile	Ser	Gly	Ser	Phe	Pro	Pro	Val	Ser	Leu	His	Tyr
				185					190					195
Ser	Tyr	Asp	Leu	Arg	Ser	Leu	Val	Ser	Gln	Leu	Phe	Lys	Arg	Asn
				200					205					210
Pro	Arg	Asp	Arg	Pro	Ser	Val	Asn	Ser	Ile	Leu	Glu	Lys	Gly	Phe
				215					220					225
Ile	Ala	Lys	Arg	Ile	Glu	Lys	Phe	Leu	Ser	Pro	Gln	Leu	Ile	Ala
				230					235					240
Glu	Glu	Phe	Cys	Leu	Lys	Thr	Phe	Ser	Lys	Phe	Gly	Ser	Gln	Pro
				245					250					255
Ile	Pro	Ala	Lys	Arg	Pro	Ala	Ser	Gly	Gln	Asn	Ser	Ile	Ser	Val
				260					265					270
Met	Pro	Ala	Gln	Lys	Ile	Thr	Lys	Pro	Ala	Ala	Lys	Tyr	Gly	Ile
				275					280					285
Pro	Leu	Ala	Tyr	Lys	Lys	Tyr	Gly	Asp	Lys	Lys	Leu	His	Glu	Lys
				290					295					300
Lys	Pro	Leu	Gln	Lys	His	Lys	Gln	Ala	His	Gln	Thr	Pro	Glu	Lys
				305					310					315
Arg	Val	Asn	Thr	Gly	Glu	Glu	Arg	Arg	Lys	Ile	Ser	Glu	Glu	Ala
				320					325					330
Ala	Arg	Lys	Arg	Arg	Leu	Glu	Phe	Ile	Glu	Lys	Asp	Lys	Glu	Arg
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Phe	Lys	Phe	Cys	Pro	Tyr	Cys	Gly	Asn	Ser	Leu	Pro	Val	Glu	Glu
				20					25					30
His	Val	Gly	Ser	Gln	Thr	Phe	Val	Asn	Pro	His	Val	Ser	Ser	Phe
				35					40					45
Gln	Gly	Ser	Gly	Ser	Arg	Pro	Pro	Thr	Pro	Lys	Ser	Ser	Pro	Gln
				50					55					60
Lys	Thr	Arg	Lys	Ser	Pro	Gln	Val	Thr	Arg	Gly	Ser	Pro	Gln	Lys
				65					70					75
Thr	Ser	Cys	Ser	Pro	Gln	Lys	Thr	Arg	Gln	Ser	Pro	Gln	Thr	Leu
				80					85					90
Lys	Arg	Ser	Arg	Val	Thr	Thr	Ser	Leu	Glu	Ala	Leu	Pro	Thr	Gly
				95					100					105
Thr	Val	Leu	Thr	Asp	Lys	Ser	Gly	Arg	Gln	Trp	Lys	Leu	Lys	Ser
				110					115					120
Phe	Gln	Thr	Arg	Asp	Asn	Gln	Gly	Ile	Leu	Tyr	Glu	Ala	Ala	Pro
				125					130					135
Thr	Ser	Thr	Leu	Thr	Cys	Asp	Ser	Gly	Pro	Gln	Lys	Gln	Lys	Phe
				140					145					150
Ser	Leu	Lys	Leu	Asp	Ala	Lys	Asp	Gly	Arg	Leu	Phe	Asn	Glu	Gln
				155					160					165
Asn	Phe	Phe	Gln	Arg	Ala	Ala	Lys	Pro	Leu	Gln	Val	Asn	Lys	Trp
				170					175					180
Lys	Lys	Leu	Tyr	Ser	Thr	Pro	Leu	Leu	Ala	Ile	Pro	Thr	Cys	Met
				185					190					195
Gly	Phe	Gly	Val	His	Gln	Asp	Lys	Tyr	Arg	Phe	Leu	Val	Leu	Pro
				200					205					210
Ser	Leu	Gly	Arg	Ser	Leu	Gln	Ser	Ala	Leu	Asp	Val	Ser	Pro	Lys
				215					220					225
His	Val	Leu	Ser	Glu	Arg	Ser	Val	Leu	Gln	Val	Ala	Cys	Arg	Leu
				230					235					240
Leu	Asp	Ala	Leu	Glu	Phe	Leu	His	Glu	Asn	Glu	Tyr	Val	His	Gly
				245					250					255
Asn	Val	Thr	Ala	Glu	Asn	Ile	Phe	Val	Asp	Pro	Glu	Asp	Gln	Ser
				260					265					270
Gln	Val	Thr	Leu	Ala	Gly	Tyr	Gly	Phe	Ala	Phe	Arg	Tyr	Cys	Pro
				275					280					285
Ser	Gly	Lys	His	Val	Ala	Tyr	Val	Glu	Gly	Ser	Arg	Ser	Pro	His
				290					295					300
Glu	Gly	Asp	Leu	Glu	Phe	Ile	Ser	Met	Asp	Leu	His	Lys	Gly	Cys
				305					310					315
Gly	Pro	Ser	Arg	Arg	Ser	Asp	Leu	Gln	Ser	Leu	Gly	Tyr	Cys	Met
				320					325					330
Leu	Lys	Trp	Leu	Tyr	Gly	Phe	Leu	Pro	Trp	Thr	Asn	Cys	Leu	Pro
				335					340					345
Asn	Thr	Glu	Asp	Ile	Met	Lys	Gln	Lys	Gln	Lys	Phe	Val	Asp	Lys
				350					355					360
Pro	Gly	Pro	Phe	Val	Gly	Pro	Cys	Gly	His	Trp	Ile	Arg	Pro	Ser
				365					370					375
Glu	Thr	Leu	Gln	Lys	Tyr	Leu	Lys	Val	Val	Met	Ala	Leu	Thr	Tyr
				380					385					390
Glu	Glu	Lys	Pro	Pro	Tyr	Ala	Met	Leu	Arg	Asn	Asn	Leu	Glu	Ala
				395					400					405
Leu	Leu	Gln	Asp	Leu	Arg	Val	Ser	Pro	Tyr	Asp	Pro	Ile	Gly	Leu
				410					415					420
Pro	Met	Val	Pro											

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PF-0614 US

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<213> Homo sapiens

<220> -

<223> 2605059

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Asp	Leu	Leu	Gly	Gln	Phe	Leu	Leu	Tyr	Pro	Pro	His	Gln	Arg	Ile
				20					25					30
Ala	Ala	Ser	Lys	Ala	Leu	Leu	His	Gln	Tyr	Phe	Phe	Thr	Ala	Pro
				35					40					45
Leu	Pro	Ala	His	Pro	Ser	Glu	Leu	Pro	Ile	Pro	Gln	Arg	Leu	Gly
				50					55					60
Gly	Pro	Ala	Pro	Lys	Ala	His	Pro	Gly	Pro	Pro	His	Ile	His	Asp
				65					70					75
Phe	His	Val	Asp	Arg	Pro	Leu	Glu	Glu	Ser	Leu	Leu	Asn	Ser	Glu
				80					85					90
Leu	Ile	Arg	Pro	Phe	Ile	Leu	Glu	Gly						
				95										

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<212> PRT

<213> Homo sapiens

<220> -

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Met	Trp	Val	Val	Pro	Pro	Ile	Gly	Ala	Glu	Phe	Leu	Gly	Thr	Glu
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Lys	Gly	Gly	Leu	Arg	Asp	Gln	Lys	Thr	Pro	Asp	Asp	His	Glu	Ala
				20					25					30
Glu	Thr	Gly	Ile	Lys	Ser	Lys	Glu	Ala	Arg	Lys	Tyr	Ile	Phe	Asn
				35					40					45
Cys	Leu	Asp	Ala	Cys	Val	Gln	Val	Asn	Met	Thr	Thr	Asp	Leu	Glu
				50					55					60
Gly	Ser	Asp	Met	Leu	Val	Glu	Lys	Ala	Asp	Arg	Arg	Glu	Phe	Ile
				65					70					75
Asp	Leu	Leu	Lys	Lys	Met	Leu	Thr	Ile	Asp	Ala	Asp	Lys	Arg	Ile
				80					85					90
Thr	Pro	Ile	Glu	Thr	Leu	Asn	His	Pro	Phe	Val	Thr	Met	Thr	His
				95					100					105
Leu	Leu	Asp	Phe	Pro	His	Ser	Thr	His	Val	Lys	Ser	Cys	Phe	Gln
				110					115					120
Asn	Met	Glu	Ile	Cys	Lys	Arg	Arg	Val	Asn	Met	Tyr	Asp	Thr	Val
				125					130					135
Asn	Gln	Ser												

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<400> 10

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gctgggtggc	tgggtgacg	gtaattcccc	gggggggtca	agtgccccaa	actgctcttg	120
gtgaaaggat	gctgtcttcc	ccgaatggcc	acttccgcct	gccttagctt	gggtgagag	180
gggacagaga	gcacctgag	gcgggccggc	caggtcttcc	cactccta	ggagctgtgg	240
ggagtggggc	cacaggcggg	gaggcagggg	gagtagtgag	tagctggtgc	caaggggccc	300
tggcgccaca	ttctggtgtc	catggggagcc	ctggggcccc	gagaggcctc	ttccctggcg	360
gctgtgcagg	gaaacctcca	cttcatgctg	actggggcgg	gcgacaggaa	ccctgggggtg	420
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1721

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<213> Homo sapiens

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g

[illegible]